

Sequence Listing

<110> Sidhu, Sachdev Weiss, Gregory A. Wells, James A.

<120> IMPROVEMENTS IN PHAGE DISPLAY

<130> P1581R2

<140> US 09/380,447 <141> 1999-09-01

<150> US 60/134,870

<151> 1999-05-19

<150> US 60/133,296

<151> 1999-05-10

<150> US 60/103,514

<151> 1998-10-08

<150> US 60/094,291

<151> 1998-07-27

<150> PCT/USUS99/16596

<151> 1999-07-22

<160> 287

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<223> Synthetic coat protein

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<221> unsure

<222> 12-30

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Met Ser Lys Ser Thr Phe Lys Lys Phe Leu Lys Xaa Xaa Xaa Xaa 10

Glu Thr Ala Ser Ala Gln Leu Ser Asn Phe Ala Ala Lys Ala Pro

Asp Asp Gly Glu Ala

<210> 2

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Way .

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<223> coat protein VIII
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 Ala Ser Ala Thr Glu Tyr Ile Gly Tyr Ala Trp Ala Met Val Val
 Val Ile Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe
 Thr Ser Lys Ala Ser
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 Val Ile Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe
 Thr Ser Lys Ala Ser
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<223> coat protein VIII

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Leu Val Val Gly Ala Thr Val Gly Ile Lys Leu Phe Lys Lys Phe

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 Thr Gln Ala Thr Asp Leu Ile Asp Gln Thr Trp Pro Val Val Thr
 Ser Val Ala Val Ala Gly Leu Ala Ile Arg Leu Phe Lys Lys Phe
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 Ser Ser Lys Ala Val
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Thr Gln Ala Ile Asp Leu Ile Ser Gln Thr Trp Pro Val Val Thr
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 taactccctg caagcc 66
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<211> 66
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<222> 19, 22, 25, 28, 31, 34, 37-38, 40-41, 43-44
<223> unknown base
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 gtttaagaaa ttcacc 66
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<222> 19-20, 22-23, 31-32, 34-35, 37-38, 43-44, 46-47
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tatcggttat gcgtgg 66
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ccgctct 57
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<223> mutagenic oligonucleotide

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ggtggaggat ccgggagctg atgagccgag ggtgacgatc cc 42
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<211> 50
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<223> P12-1 variant
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 Tyr Met Leu Leu Val Glu Ala Ser Pro Trp Ala Ala Lys Ala Pro
 Asp Asp Gly Glu Ala
<210> 34
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cvvcvvcvvc vvcvvcvvcg gcggtgccga gggtgacgat ccc 93
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<220>

<223> oligonucleotide linker library

<400> 35

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vcvvcvvcvv cvvcvvcvvc vvcvvcvvcg ccgagggtga cgatccc 97
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<223> oligonucleotide linker library
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 tatcggttat gcgtgg 66
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<211> 66
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 cattgtcggc gcaact 66
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gcaaaagcgg cctataacgc tcttgaggat att 33
<210> 43
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<400> 43
tatgaggctc ttgaggccat tgctactaac tat 33
<210> 44
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 tatcggttat gcgtgg 66
<210> 46
<211> 66
<212> DNA
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 cgagggtgac gatccc 66
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<211> 60
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 cggttatgcg 60
<210> 48
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cattgtcggc gcaact 66
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agctccagcg ataagggtga gaagttcgct aga 33
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 cggttatgcg 60
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tgttgat 57
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 Gln Tyr Gly Thr Pro Asp Thr Asp
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 accatcacca tcaccatgcg 120
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 Ala Gln Leu Ser Asn Phe Ala Ala Lys Ala Pro Asp Asp Gly Glu
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Asp Asp Gly Glu Ala Ala Ala Asp Ala

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- <400> 124
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- tgatgttgat aataattgga tttgggctgt cggtattatt tacatgctcc 100

tcgtggaggc gtcgccctgg gctgctaagg cgccagacga tggtgaagct 150

- <210> 125
- <211> 48
- <212> DNA
- <213> Artificial sequence
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- <223> mutagenic oligonucleotide library
- <220>

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 t 51
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<211> 30
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 Ile Lys Asp Glu Gly Phe Ala Arg Ala Ala
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